Sequences removed

---------- Forwarded message ---------
From: <gb-admin@ncbi.nlm.nih.gov>
Date: Tue, Apr 28, 2020 at 11:53 AM
Subject: GenBank MT063207-MT063722
To: <aashaffer@ucdavis.edu>, <tgoldstein@ucdavis.edu>

Dear Dr. Gibson:

Regarding your request:

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Subject: Remove accessions from Genbank
Date: Tue, 28 Apr 2020 16:51:58 +0000

We are requesting the removal of 728 sequences from GenBank. Please find the
list attached as a text file. Thank you!
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You will be contacted about the remaining records separately.

We have temporarily delayed the release of your GenBank records
MT063207-MT063722 to get clarification about your request.

What is the reason for deleting MT063224-MT063227, MT063351-MT063358,
MT063391-MT063392, MT063603-MT063610, MT063621-MT063631, MT063642,
MT063687, MT063689-MT063692, MT063710-MT063711, MT063713?

If the issue can be resolved by updating the current records,
send the update in the following formats so that the new information
can be added accurately.

If you are updating multiple records, please send a list of all
accessions to be updated at the top of your request.

[1] Publication (Reference) Updates:

    [a] If the PMID or DOI are publicly available please send the
       information as a tab-delimited table as follows:

acc. num.	PMID
MHXXXX01	29980901
MHXXXX02	29980901

or

acc. num.	DOI

```
MHXXXX01      10.1000/xyz123
MHXXXX02      https://doi.org/10.1000/xyz123doi
```

[b] For all other updates, please provide the revised information
in a tab-delimited table. You must replace any non-ASCII
characters (for example, characters with accents and umlauts)
with the appropriate English letters.

The complete list of revised author names should be provided in the
following format: first_initial middle_initial surname, etc., For example:

```
acc. num.        authors          title
MHXXXX01         J. A. Smith    Identification of gene A
MHXXXX02         X. P. Weng, J. Doe   Identification of gene B
```

A list of publication fields in the format to be used in the column headers
is available at:

https://www.ncbi.nlm.nih.gov/Genbank/update.html

[2] Source (Organism) Updates

Send the update as a tab-delimited table, like this:

```
acc. num.     organism        strain
MHXXXX01      Escherichia coli      82
MHXXXX02      Bacillus subtilis     ABC
```

A list of accepted GenBank source qualifiers can be found at
https://www.ncbi.nlm.nih.gov/Sequin/modifiers.html

[3] Sequence Updates

Send the complete new sequences in a single fasta file. If the accession
number has been assigned, this number must be used as the identifier,
for example:

```
>MHXXXX01
cggtaataatggaccttggaccccggcaaagcggagagac
>MHXXXX02
ggaccttggaccccggcaaagcggagagaccggtaataat
```

[4] Feature (Annotation) Updates

If you are adding annotation or changing locations of features then send us
the features in a tab-delimited, 5-column table. The table format is described
at https://www.ncbi.nlm.nih.gov/Sequin/table.html.

  a. If the record is publicly released and has annotation, you can
      download the existing annotation .tbl file by retrieving the
      record and clicking on the 'Send to' option. Choose 'File' as
      Destination and then Format 'Feature table'. Edit this table
      and send to us via email.

b. If the record is not yet publicly released, let us know, and we
   will send you a 5-column table with the current annotation for
   you to edit and return to us.

Please maintain the tab structure of the table when editing.

For further information about other methods of updating your current
GenBank records, see: https://www.ncbi.nlm.nih.gov/Genbank/update.html

Please send your reply to: gb-admin@ncbi.nlm.nih.gov

Thank you for your cooperation, and we look forward to receiving your
update request.

Sincerely,

Francescopaolo Di Cello, PhD
Contractor
The GenBank Submissions Staff
Bethesda, Maryland USA
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gb-admin@ncbi.nlm.nih.gov (for replies/updates to records in GenBank)
info@ncbi.nlm.nih.gov   (for general questions regarding GenBank)
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